

# Supplementary material

Predicting plant Rubisco kinetics from RbcL sequence data using machine learning

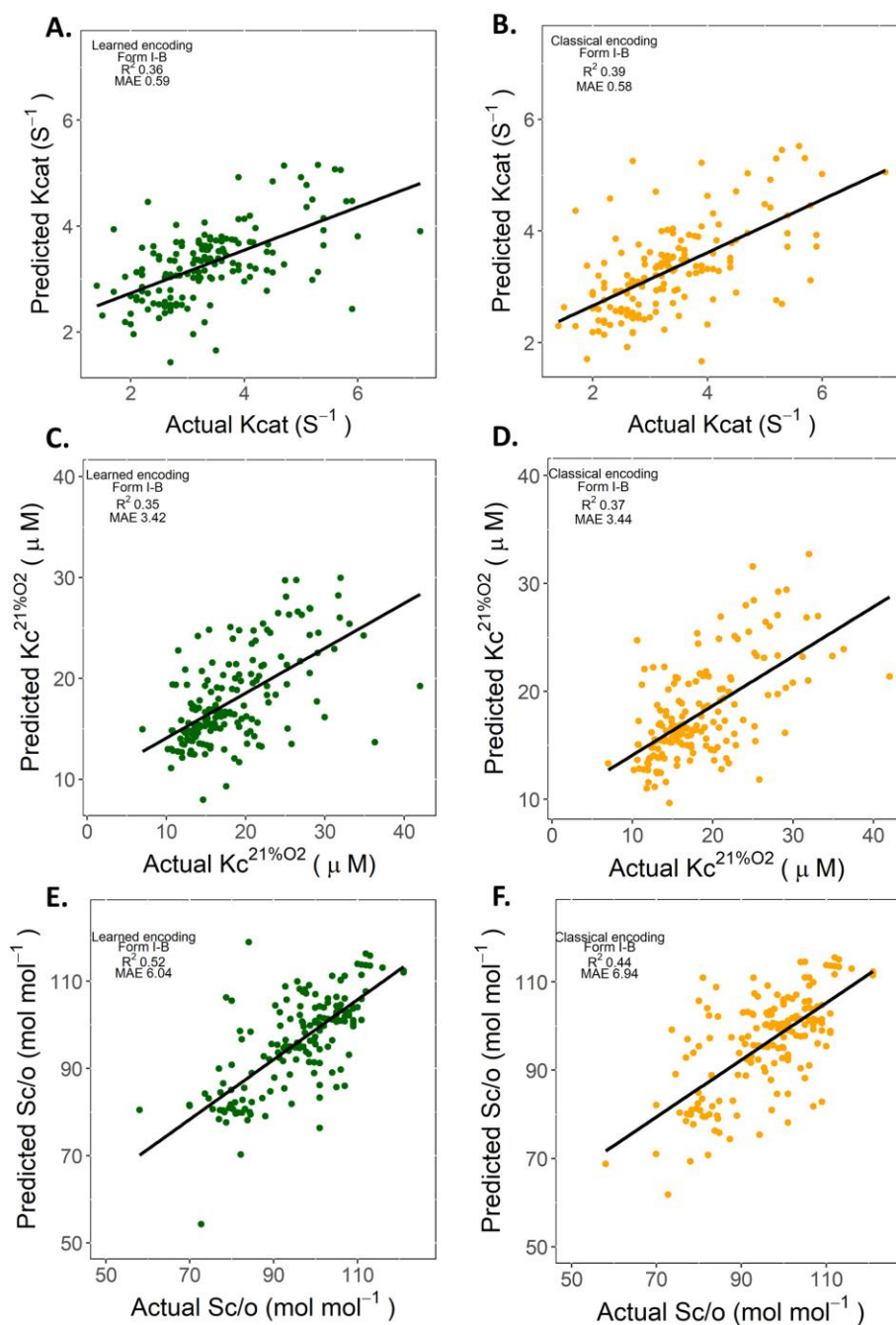
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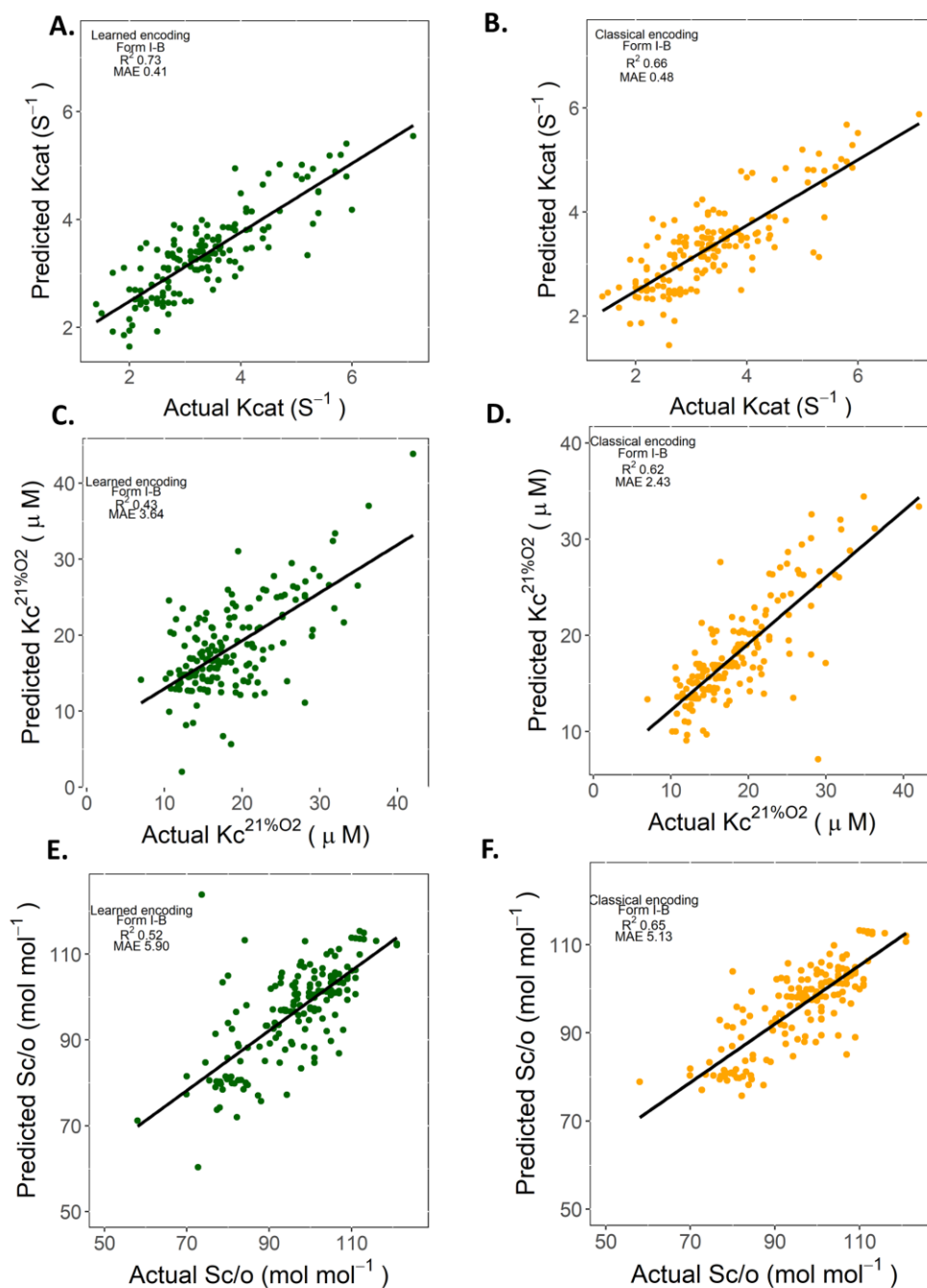
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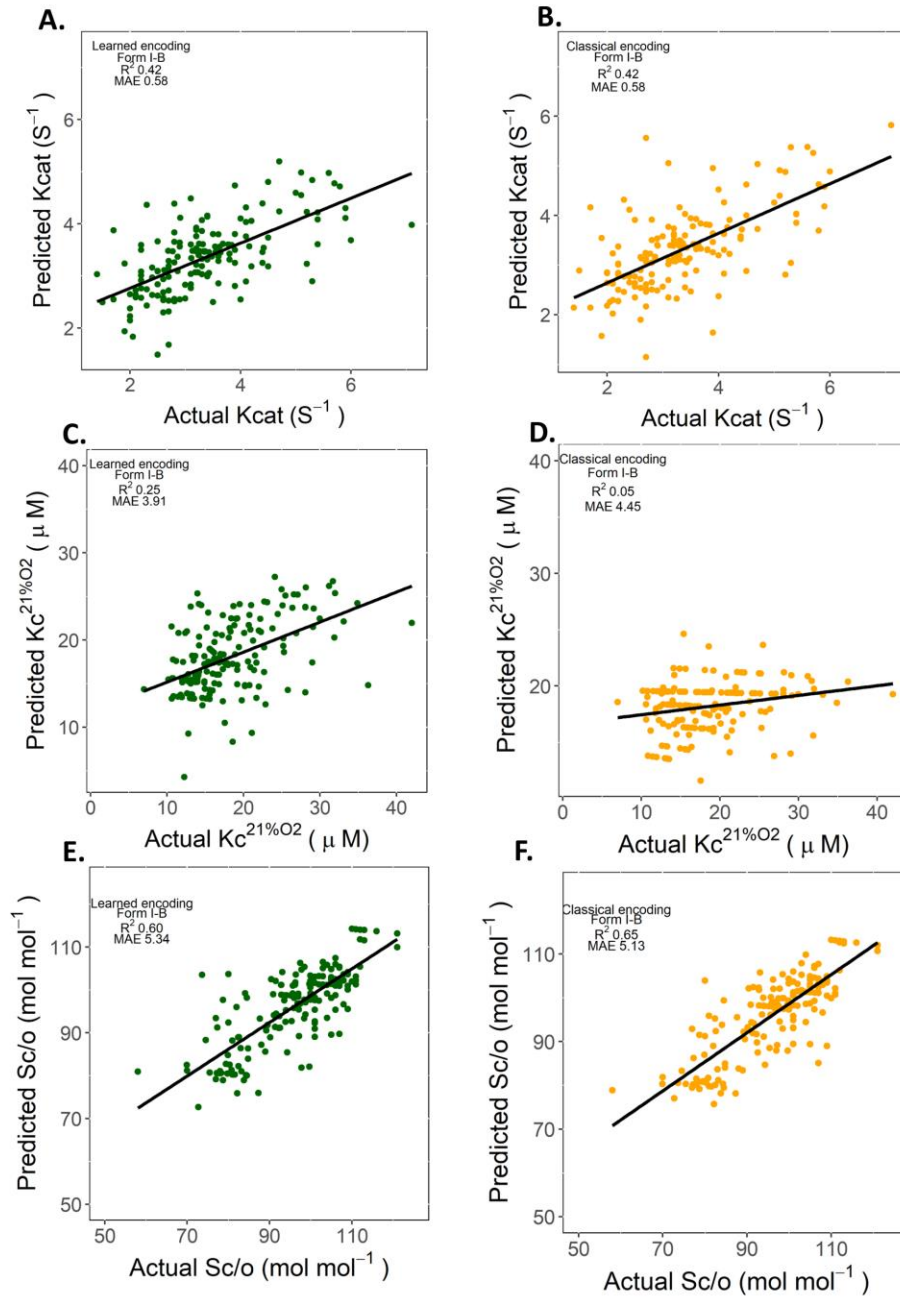
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**Figure S1.** Comparison between predicted and actual carboxylation turnover rate (Kcat : s<sup>-1</sup>), Michaelis-Menten constant for CO<sub>2</sub> at ambient O<sub>2</sub> (Kc<sup>21%O<sub>2</sub></sup>: μM) and specificity for CO<sub>2</sub> over O<sub>2</sub> (Sc/o: mol mol<sup>-1</sup>) at 25°C. Determined using leave-one-out cross-validation with the learned encodings (Rives et al., 2021) (green) and classical encodings (orange) encoded by a single Matern 5/2 kernel. The poor performance of the single Matern 5/2 kernel approach justified the use of an additive Matern 5/2 kernel for the final gaussian process models (see paper Figure 2).



**Figure S2.** Comparison between predicted and actual carboxylation turnover rate (Kcat : s<sup>-1</sup>), Michaelis-Menten constant for CO<sub>2</sub> at ambient O<sub>2</sub> (Kc<sup>21%O<sub>2</sub></sup>: μM) and specificity for CO<sub>2</sub> over O<sub>2</sub> (Sc/o: mol mol<sup>-1</sup>) at 25°C. Determined using leave-one-out cross-validation with the learned encodings (Rives et al., 2021) (green) and classical encodings (orange) encoded by an additive linear kernel. The poor performance of the additive linear kernel approach justified the use of an additive Matern 5/2 kernel for the final gaussian process models (see paper Figure 2).

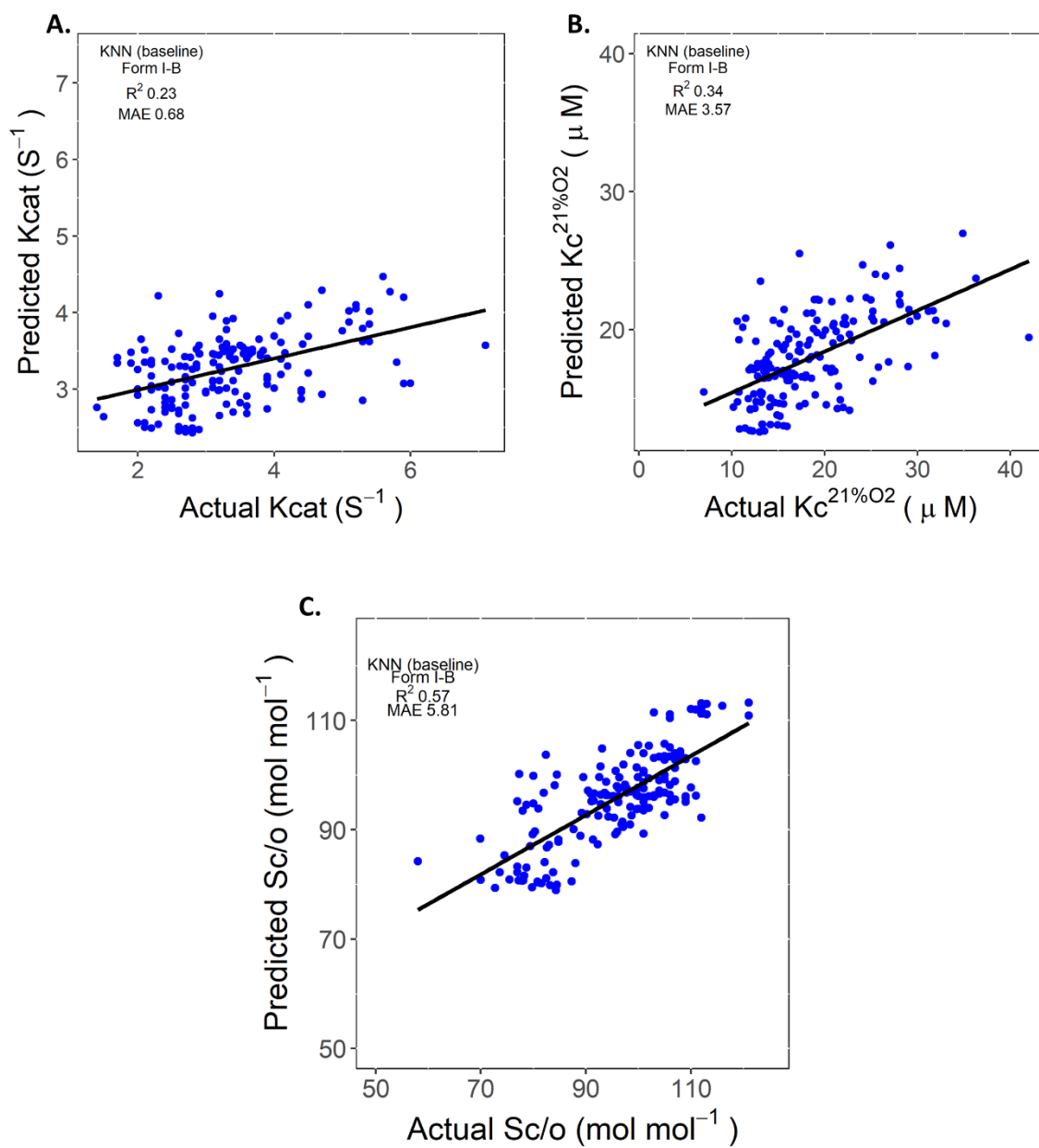


**Figure S3.** Comparison between predicted and actual carboxylation turnover rate ( $K_{cat}$  :  $s^{-1}$ ), Michaelis-Menten constant for  $CO_2$  at ambient  $O_2$  ( $K_c^{21\%O_2}$ :  $\mu M$ ) and specificity for  $CO_2$  over  $O_2$  ( $Sc/o$ :  $mol\ mol^{-1}$ ) at  $25^\circ C$ . Determined using leave-one-out cross-validation with the learned encodings (Rives et al., 2021) (green) and classical encodings (orange) encoded by a single linear kernel. The poor performance of the single linear kernel approach justified the use of an additive Matern 5/2 kernel for the final gaussian process models (see paper Figure 2).

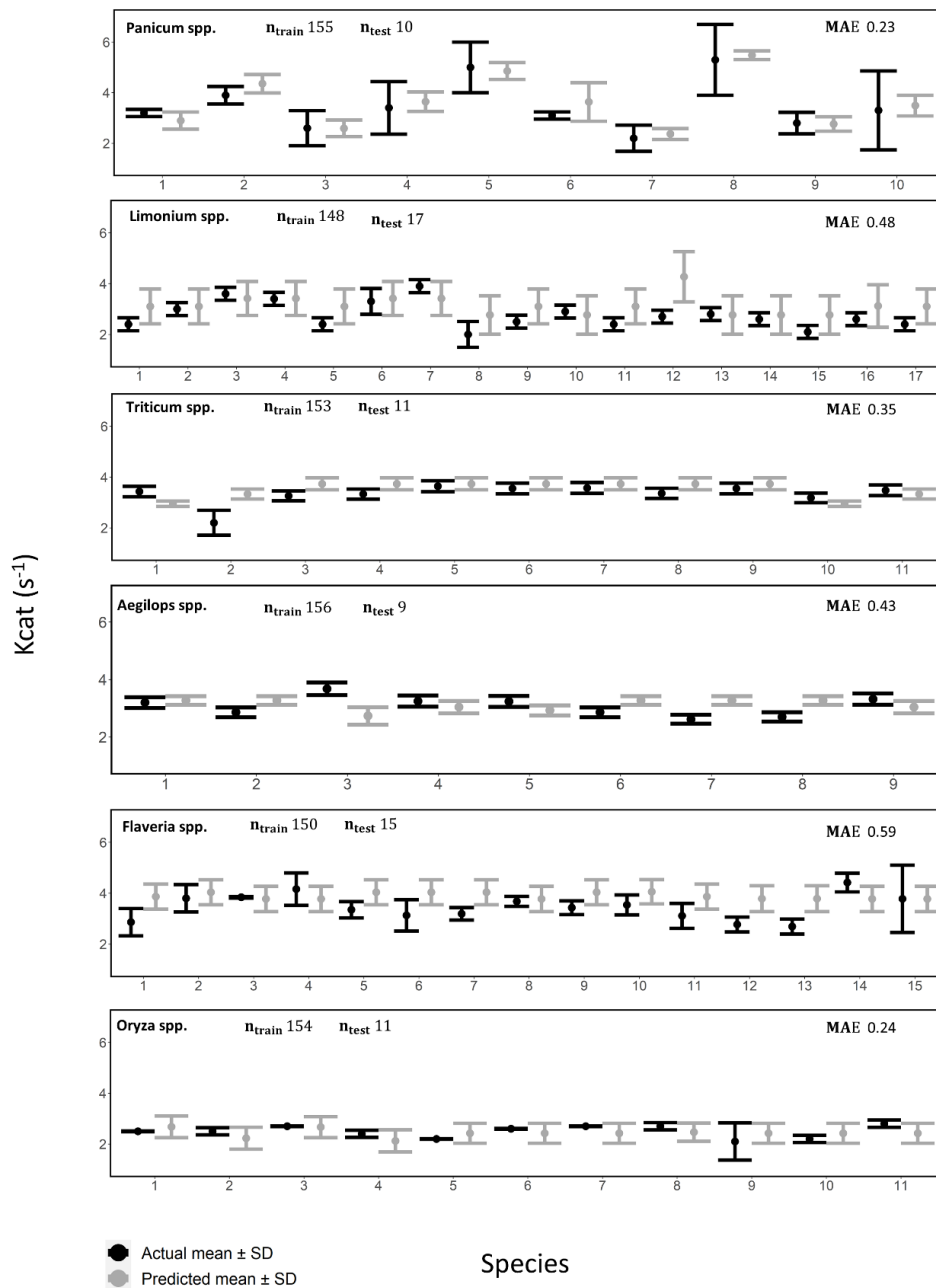
**Table S1.** Average performance of the learned encoding scheme across five randomly chosen sets of untrained weights determined using leave-one-out cross validation.

Model configuration	Task	R <sup>2</sup>	MAE
Additive Matern 5/2 kernel	Kcat	0.79 ± 0.01	0.40 ± 0.01
Additive Matern 5/2 kernel	Kc <sup>21%O2</sup>	0.82 ± 0.03	1.97 ± 0.17
Additive Matern 5/2 kernel	Sc/o	0.79 ± 0.01	4.03 ± 0.12
Single Matern 5/2 kernel	Kcat	0.30 ± 0.02	0.65 ± 0.01
Single Matern 5/2 kernel	Kc <sup>21%O2</sup>	0.25 ± 0.01	3.63 ± 0.04
Single Matern 5/2 kernel	Sc/o	0.66 ± 0.01	5.18 ± 0.06
Additive linear kernel	Kcat	0.63 ± 0.03	0.47 ± 0.02
Additive linear kernel	Kc <sup>21%O2</sup>	0.56 ± 0.06	3.16 ± 0.22
Additive linear kernel	Sc/o	0.70 ± 0.04	4.89 ± 0.32
Single linear kernel	Kcat	0.37 ± 0.02	0.60 ± 0.01
Single linear kernel	Kc <sup>21%O2</sup>	0.20 ± 0.03	3.94 ± 0.09
Single linear kernel	Sc/o	0.63 ± 0.01	5.41 ± 0.07

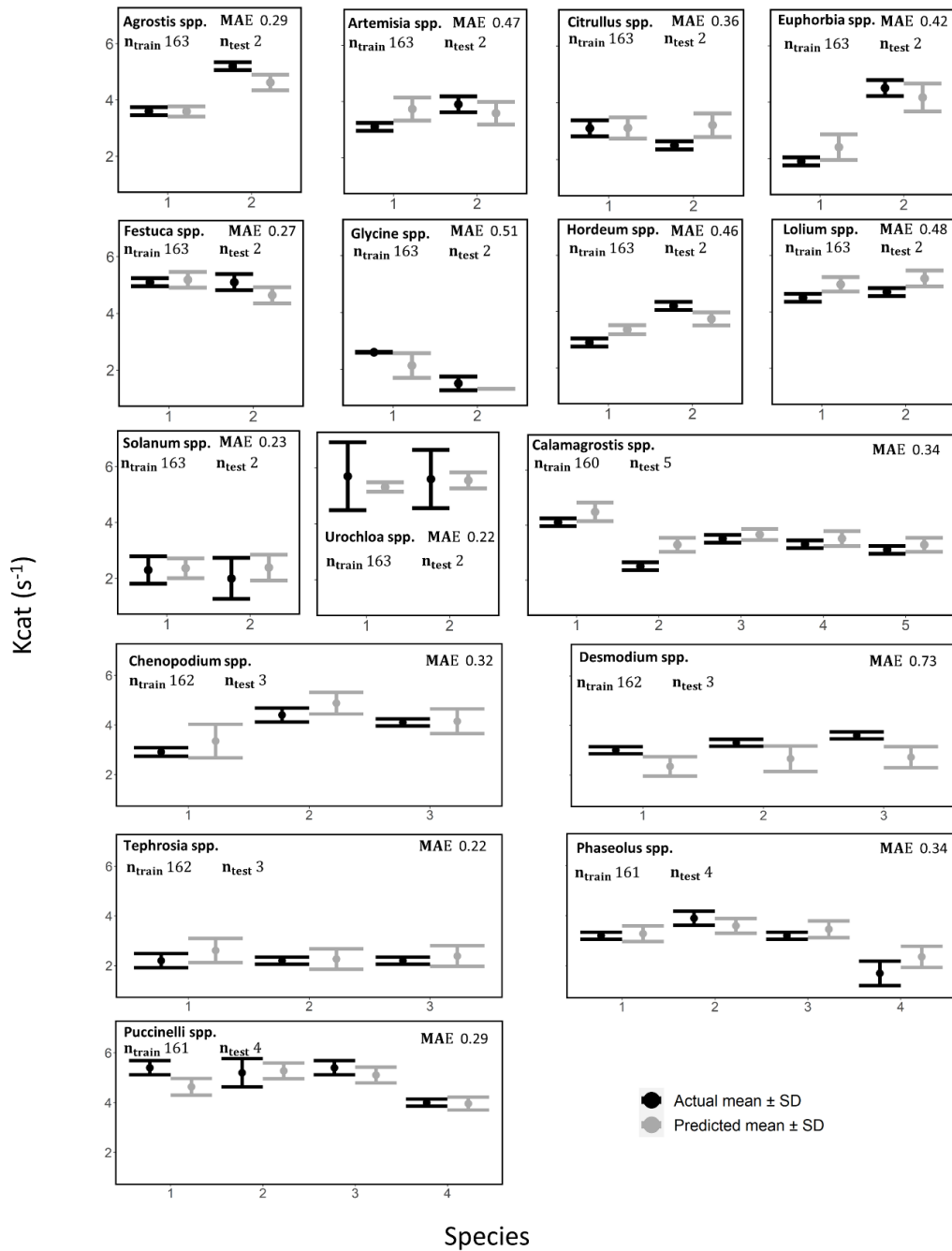
The learned encoding scheme is based on a fixed pre-trained neural network. Randomly assigning untrained weights led to minor differences in performance. This demonstrates that the boost in performance from the pretrained learned encoding scheme (e.g. Figure 2) is most likely driven by the neural network architecture. Mean ± SD performance metrics across five randomly chosen sets of untrained weights. Mean absolute errors (MAE) are in the units of the corresponding task outcomes i.e. Kcat (s<sup>-1</sup>), Kc<sup>21%O2</sup> (μM) or Sc/o (mol mol<sup>-1</sup>).



**Figure S4.** Comparison between predicted and actual carboxylation turnover rate ( $K_{cat}$  :  $s^{-1}$ ), Michaelis-Menten constant for  $CO_2$  at ambient  $O_2$  ( $K_c^{21\%O_2}$ :  $\mu M$ ) and specificity for  $CO_2$  over  $O_2$  ( $Sc/o$ :  $mol\ mol^{-1}$ ) at  $25^\circ C$ . Determined using leave-one-out cross-validation with K-nearest neighbour (KNN) sequence-similarity (baseline) models.

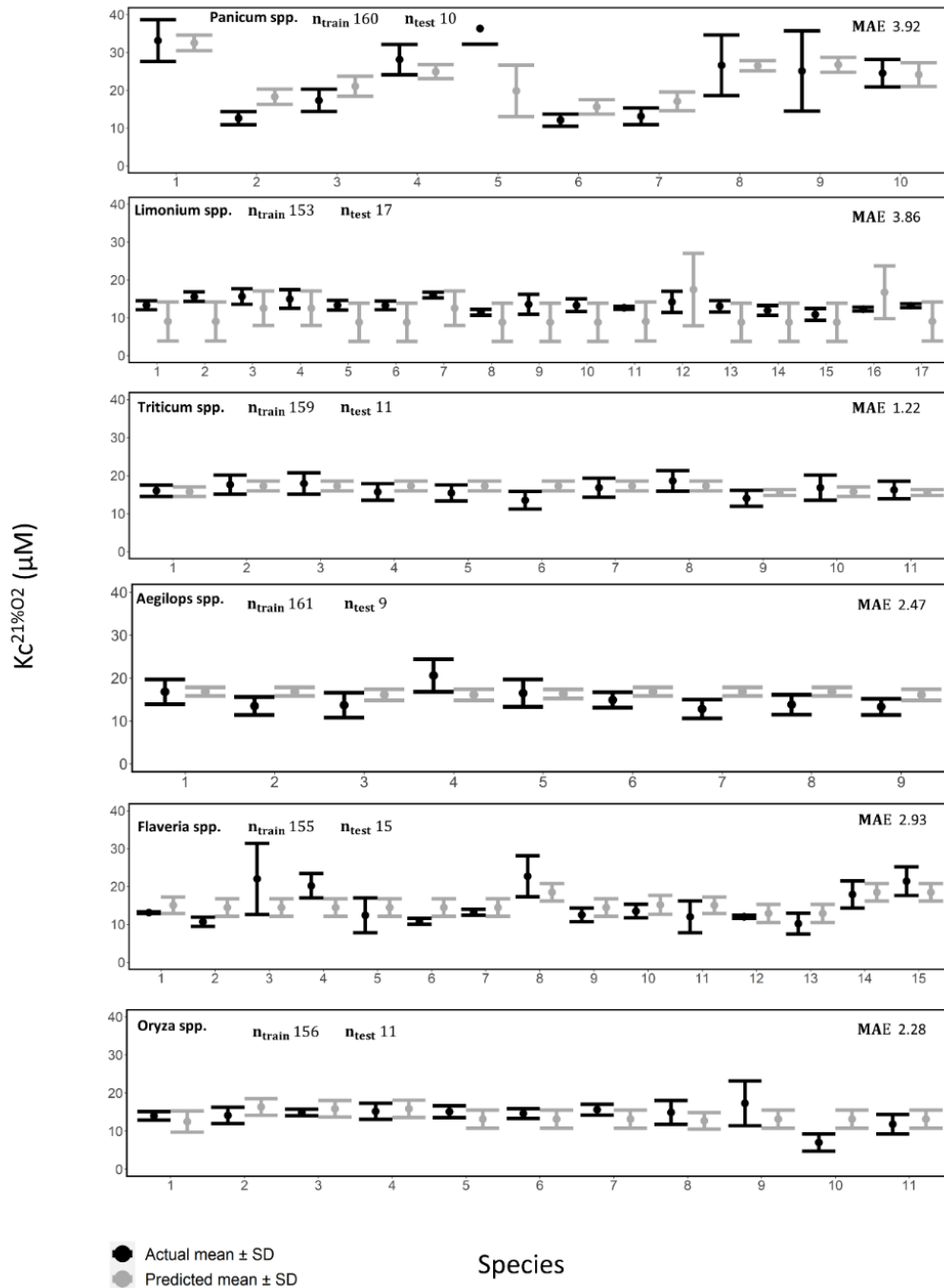


**Figure S5.** Leave-genus-out cross validation plots for carboxylation turnover rate ( $K_{cat}$ ). Another validation framework that involved leaving a genus out during model training and then evaluating the model using the unseen genus. For example, *Panicum* spp. were removed during model training leaving 155 other species for training ( $n_{train}$ ) and the model performance was assessed using the 10 unseen *Panicum* spp. ( $n_{test}$ ). This process was repeated for each genus with more than one species. Each pair of results show the actual (black) and predicted (grey) mean  $K_{cat} \pm SD$ . Actual or experimental SDs were calculated from the reported standard errors (SE) and number of replicates (see methods section). The mean absolute error (MAE) between the predicted and actual values are shown for each genus group.



**Figure S5 continued.** Leave-genus-out cross validation plots for Kcat.





**Figure S6.** Leave-genus-out cross validation plots for the Michaelis-Menten constant for  $CO_2$  at ambient atmospheric  $O_2$  ( $K_c^{21\%O_2}$ ). Same process as figure S4. Each pair of results show the actual (black) and predicted (grey) mean  $K_c^{21\%O_2} \pm SD$ . Actual or experimental SDs were calculated from the reported standard errors (SE) and number of replicates (see methods section). The mean absolute error (MAE) between the predicted and actual values are shown for each genus group.

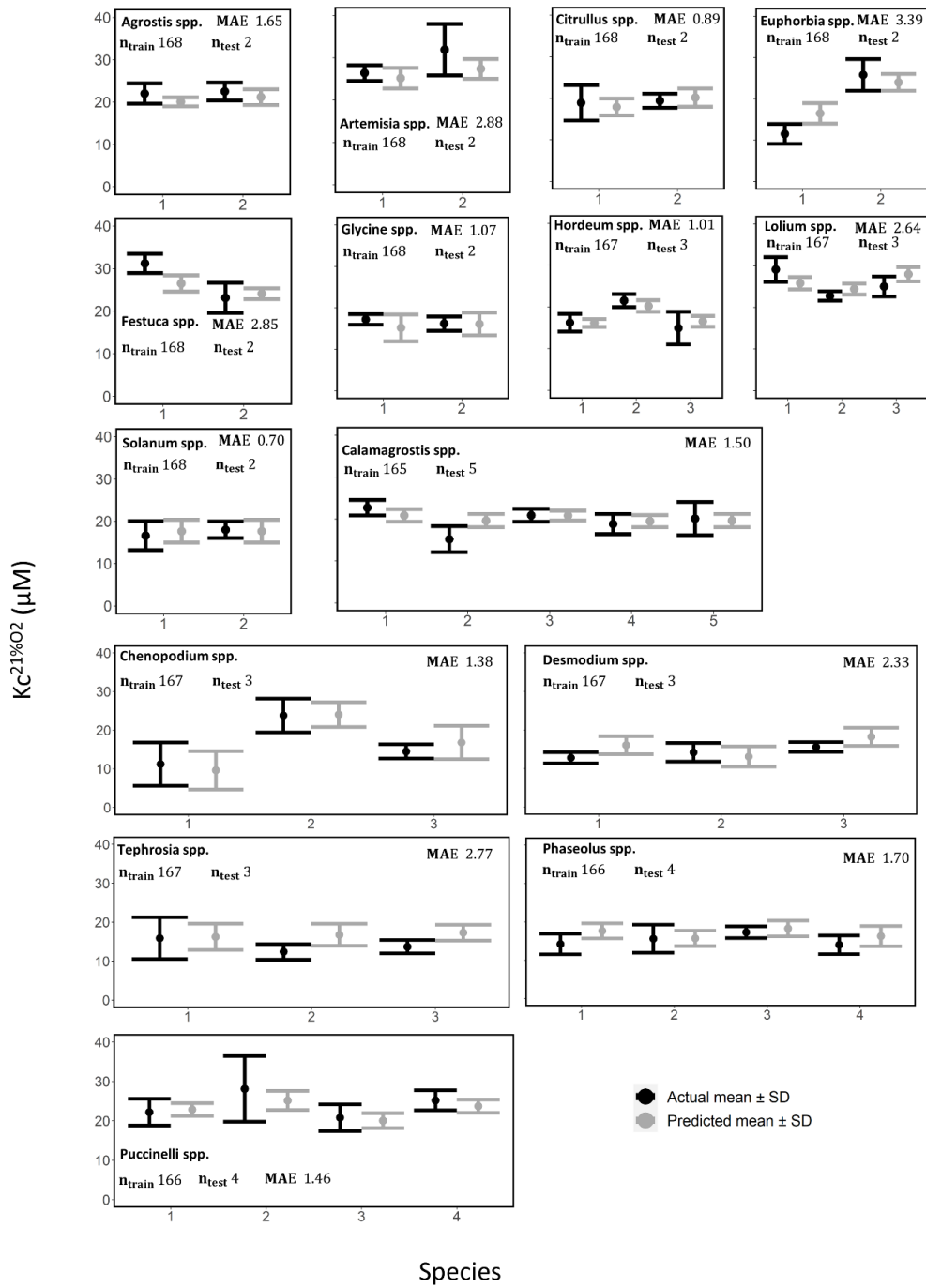
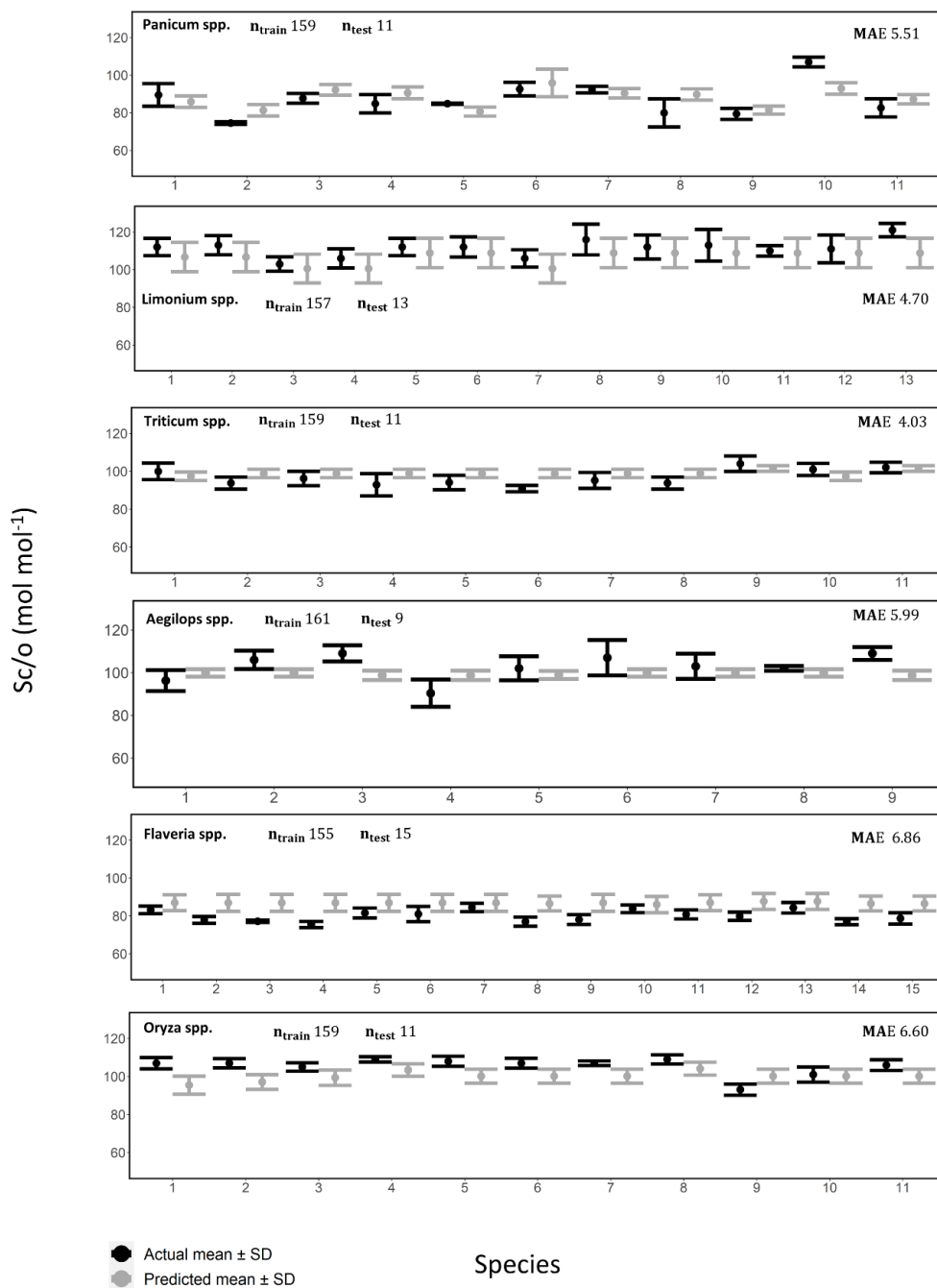


Figure S6 continued. Leave-genus-out cross validation plots for  $Kc^{21\%O_2}$ .



**Figure S7.** Leave-genus-out cross validation plots for the specificity for CO<sub>2</sub> over O<sub>2</sub> (Sc/o). Same process as figure S4. Each pair of results show the actual (black) and predicted (grey) mean Sc/o ± SD. Actual or experimental SDs were calculated from the reported standard errors (SE) and number of replicates (see methods section). The mean absolute error (MAE) between the predicted and actual values are shown for each genus group.

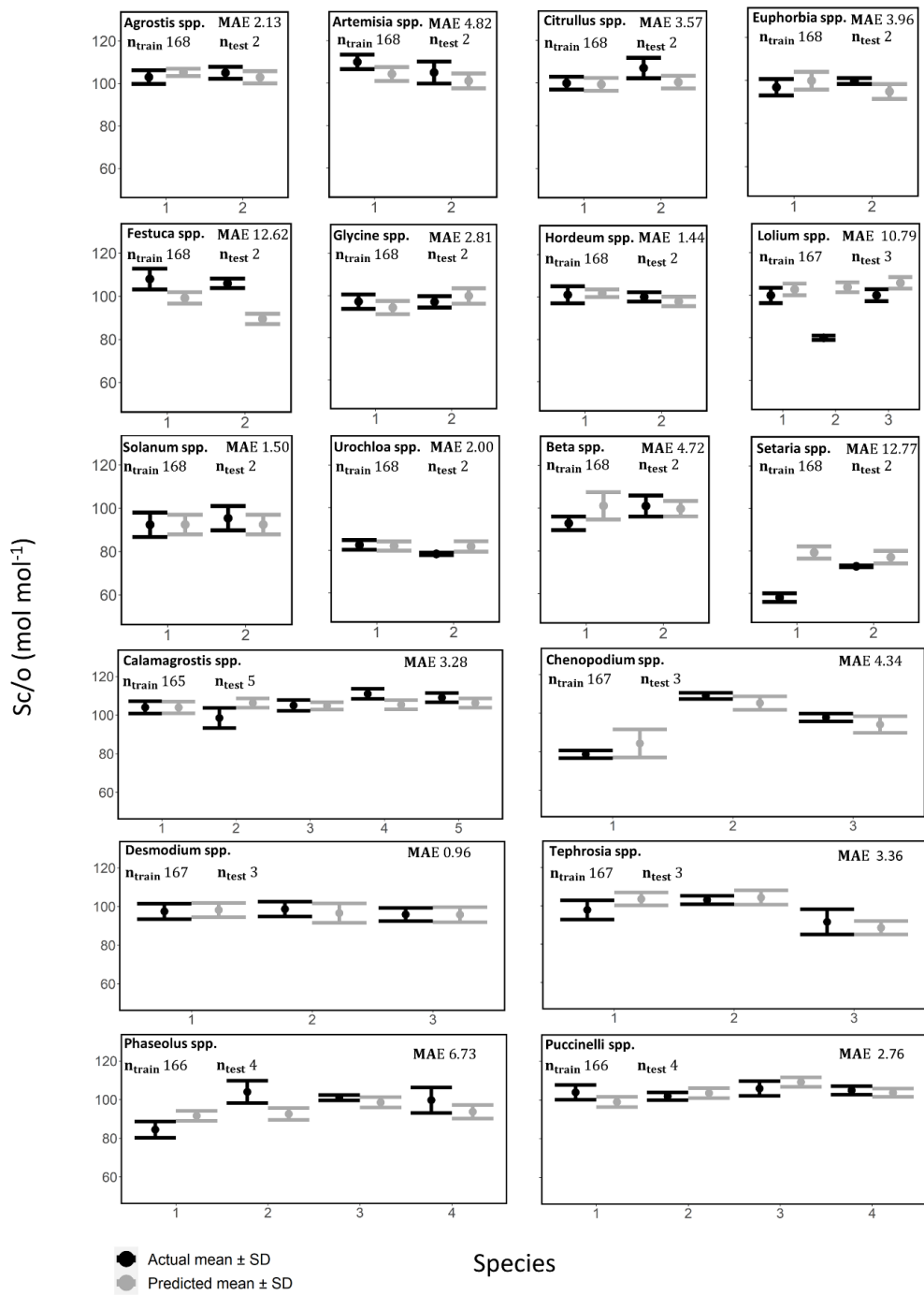
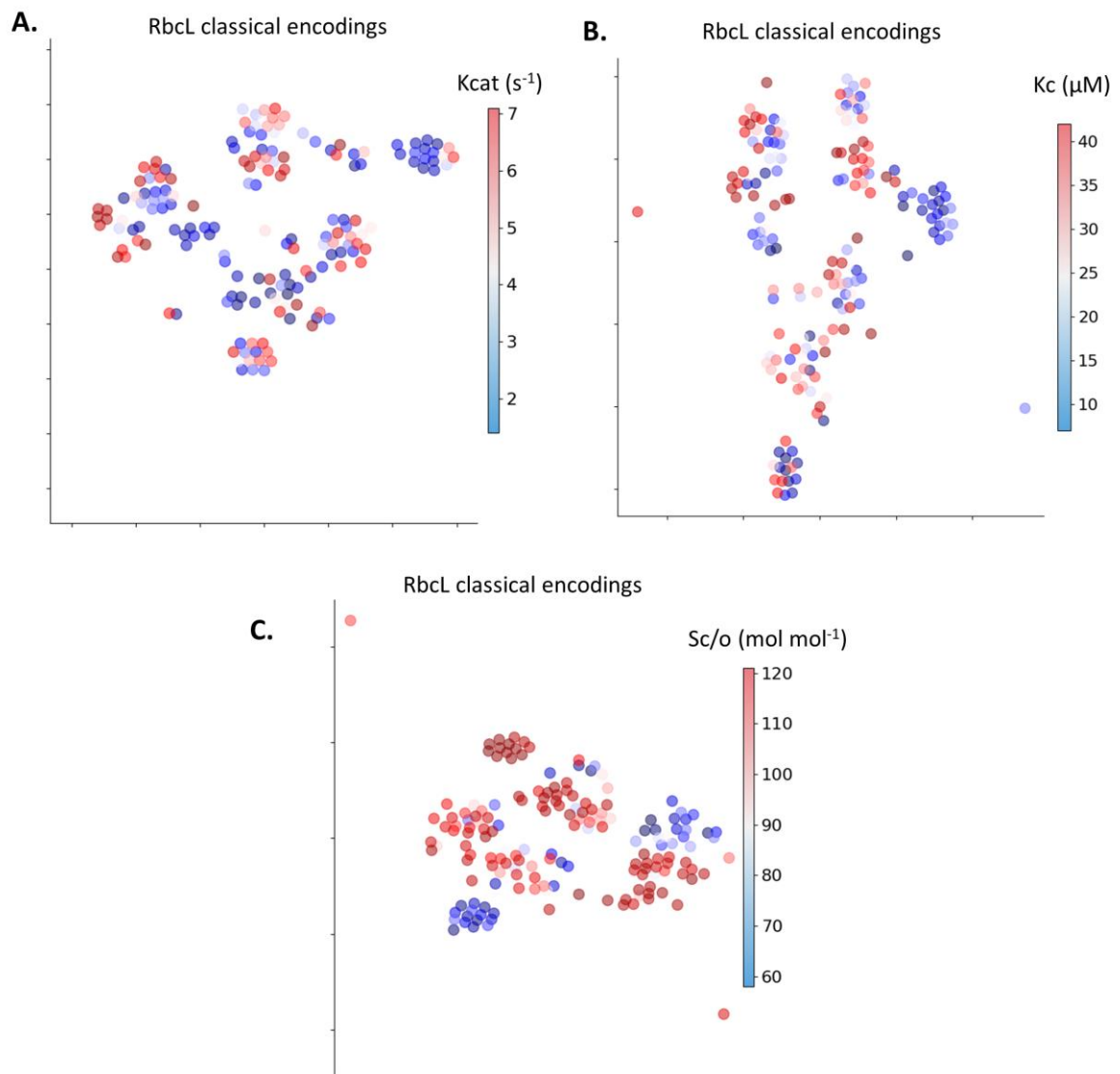
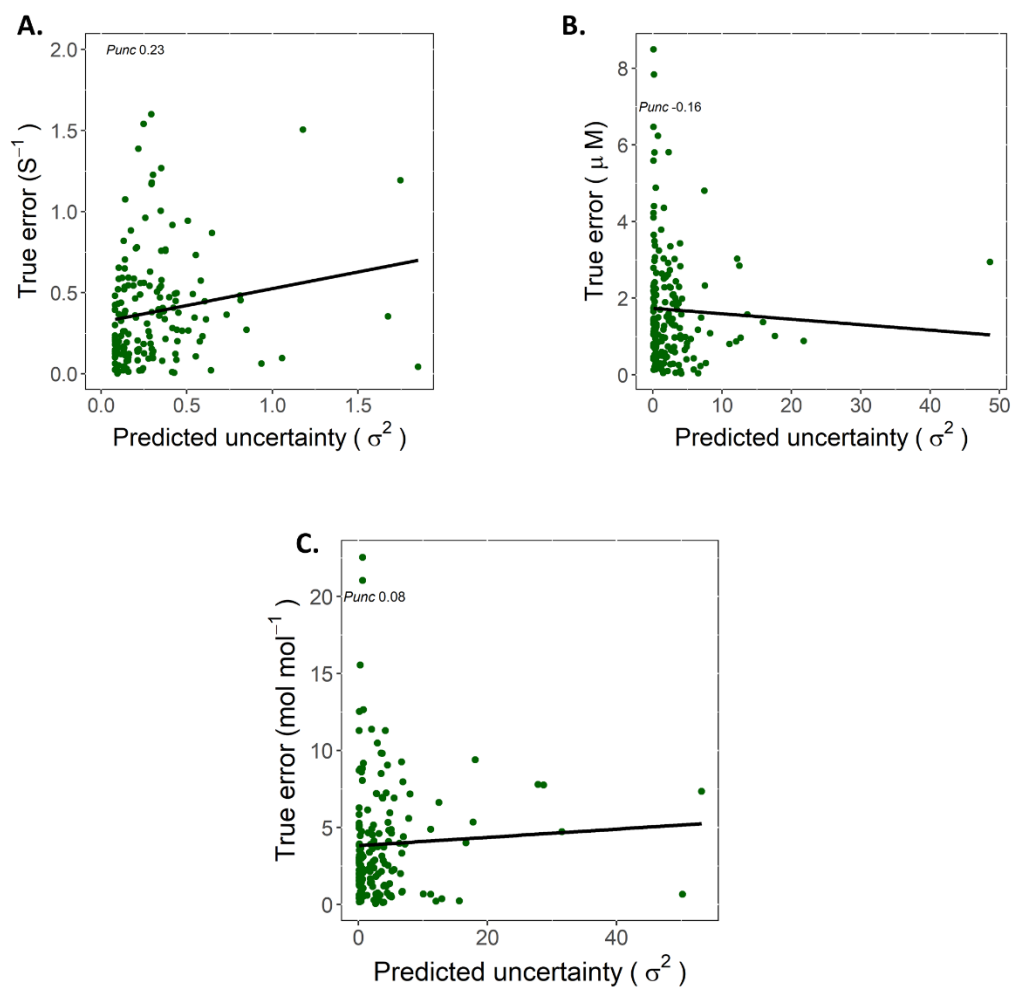


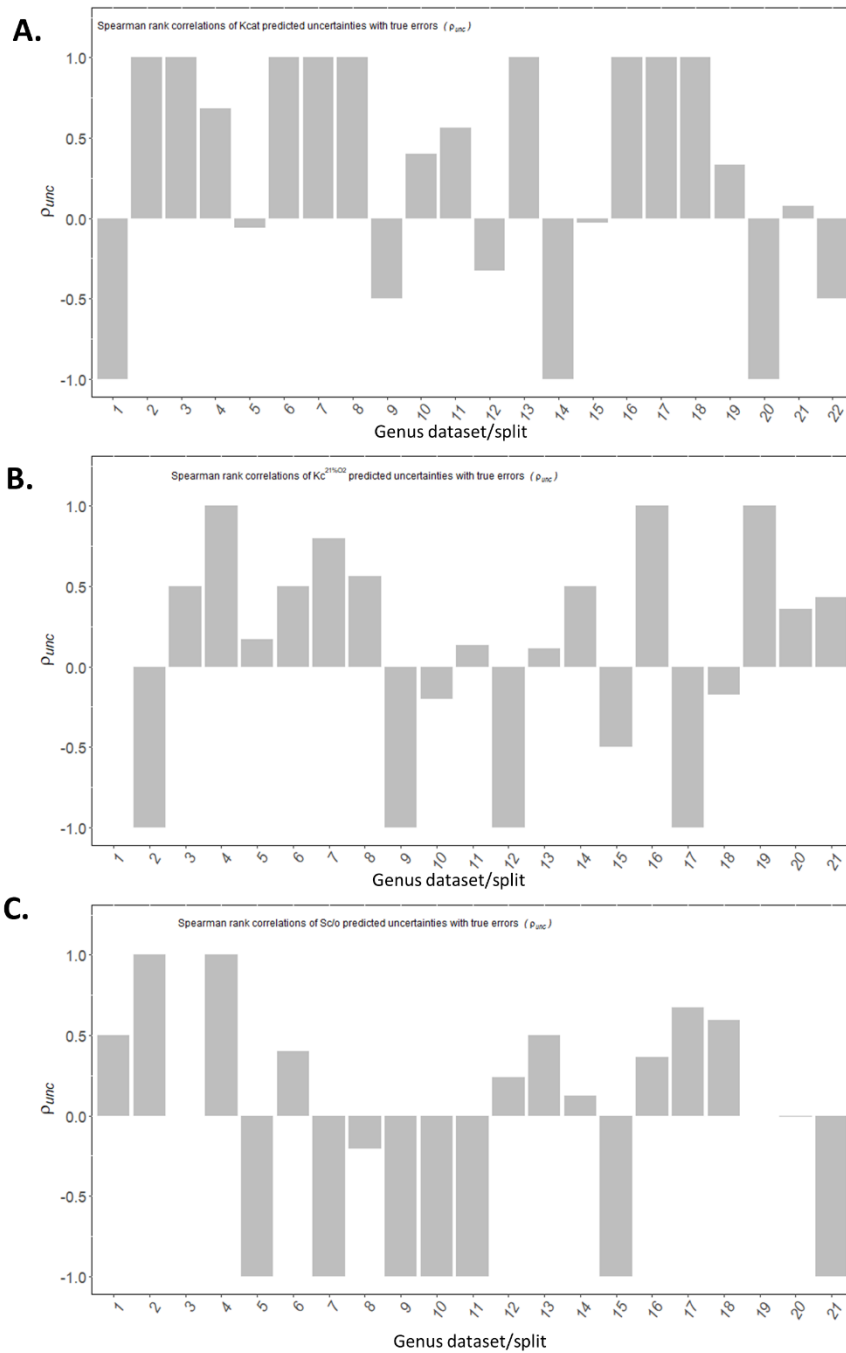
Figure S7. Continued. Leave-genus-out cross validation plots for  $Sc/o$ .



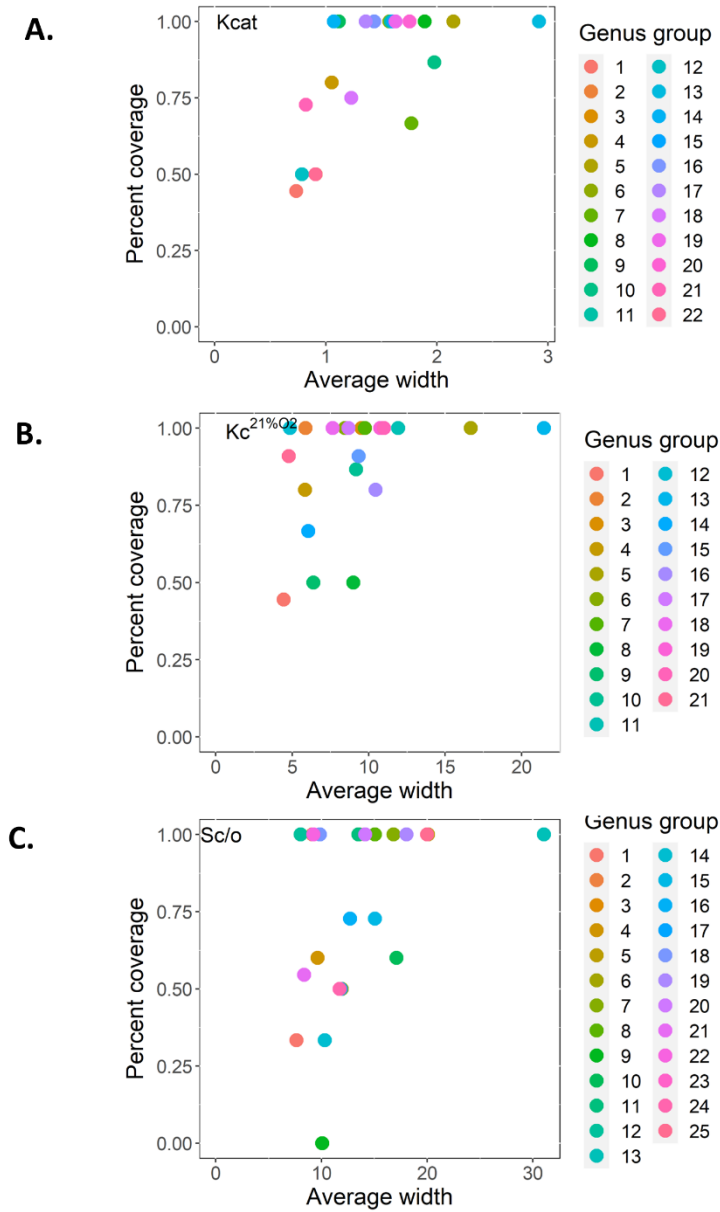
**Figure S8.** Visualization of the Rubisco large subunit (RbcL) classical encodings used in the fully trained Gaussian process (GP) models. Each data point represents an RbcL classical encoding with (A) carboxylation turnover rate ( $K_{cat}$ :  $s^{-1}$ ) ( $n=165$ ), (B) Michaelis-Menten constant for  $CO_2$  at ambient atmospheric  $O_2$  ( $K_c^{21\%O_2}$ :  $\mu M$ ) ( $n=170$ ) and (C) specificity for  $CO_2$  over  $O_2$  ( $Sc/o$ :  $mol\ mol^{-1}$ ) ( $n=170$ ).



**Figure S9.** Spearman rank correlations ( $\rho_{unc}$ ) of leave-one-out cross validation predicted uncertainties ( $\sigma^2$ ) and true errors for (A) carboxylation turnover rate ( $K_{cat} : s^{-1}$ ), (B) Michaelis-Menten constant for  $CO_2$  at ambient  $O_2$  ( $K_C^{21\%O_2} : \mu M$ ) and (C) specificity for  $CO_2$  over  $O_2$  ( $Sc/o : mol\ mol^{-1}$ ). Determined using the learned encodings (Rives et al., 2021) encoded by an additive Matern 5/2 kernel.

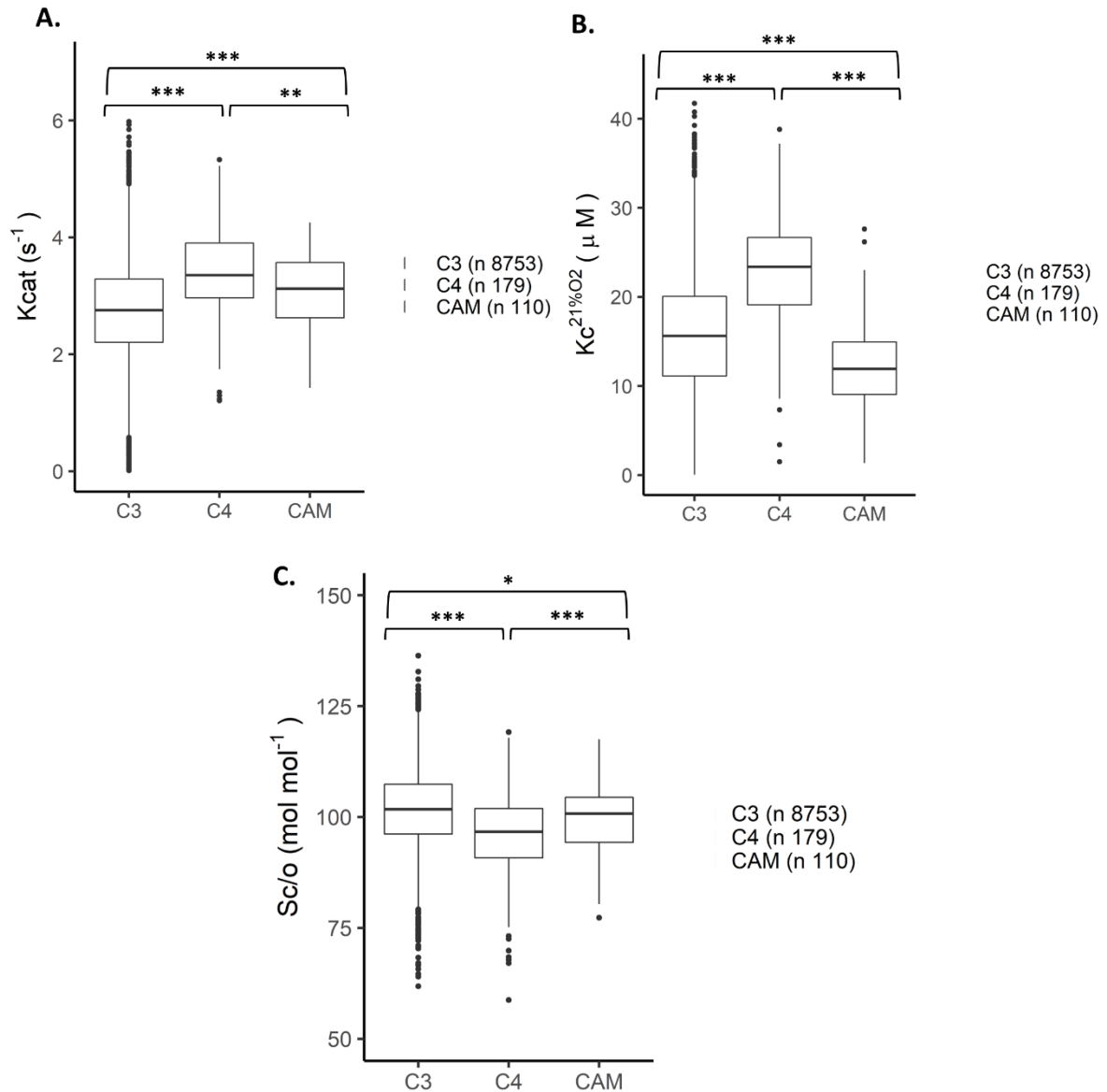


**Figure S10.** Spearman rank correlations ( $\rho_{unc}$ ) of leave-genus-out cross validation predicted uncertainties ( $\sigma^2$ ) and true errors for (A) carboxylation turnover rate (Kcat :  $s^{-1}$ ), (B) Michaelis-Menten constant for  $CO_2$  at ambient  $O_2$  ( $Kc^{21\%O_2}$ :  $\mu M$ ) and (C) specificity for  $CO_2$  over  $O_2$  (Sc/o:  $mol\ mol^{-1}$ ). Determined using the learned encodings (Rives et al., 2021) encoded by an additive Matern 5/2 kernel.  $\rho_{unc}$  values are ordered by the mean absolute error (MAE) for each genus group.

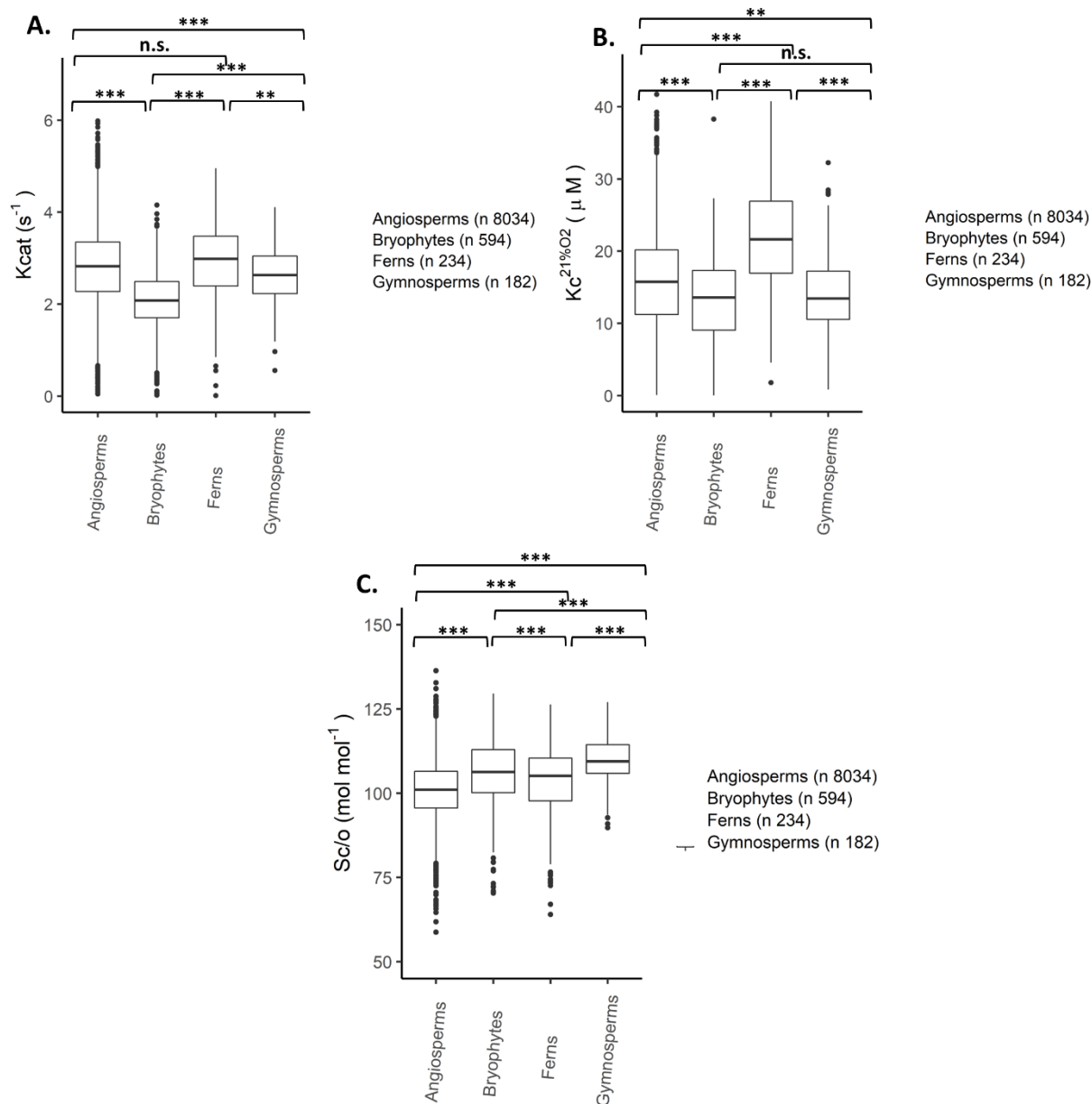


**Figure S11.** Coverage vs average width ( $4\sigma$ ) for (A) carboxylation turnover rate ( $K_{cat}$  :  $s^{-1}$ ), (B) Michaelis-Menten constant for  $CO_2$  at ambient  $O_2$  ( $Kc^{21\%O_2}$ :  $\mu M$ ) and (C) specificity for  $CO_2$  over  $O_2$  ( $Sc/o$ :  $mol\ mol^{-1}$ ) leave-genus-out cross validation predictions. Coverage is the percentage of actual mean values that fall within the predicted 95% confidence intervals ( $\pm 2\sigma$ ). High coverage and low width indicates good model uncertainty. As predicted mean values become increasingly out of distribution, we would like width to increase indicating increasing model uncertainty and coverage to remain high.





**Figure S12.** Box plots depict (A) carboxylation turnover rate ( $K_{cat}$ :  $s^{-1}$ ), (B) Michaelis-Menten constant for  $CO_2$  at ambient atmospheric  $O_2$  ( $K_c^{21\%O_2}$ :  $\mu M$ ) and (C) specificity of  $CO_2$  to  $O_2$  ( $Sc/o$ :  $mol\ mol^{-1}$ ) predictions made for the wild-type form IB (plants) Rubisco large subunit (RbL) sequence-space using the fully trained Gaussian process (GP) models with the learned encoding scheme. Predictions are shown for all known land plant Rubiscos. Predictions were grouped by photosynthesis metabolism type ( $C_3$ ,  $C_4$  or CAM). Box plot horizontal lines show the median value, and the box and whisker represent the 25<sup>th</sup> and 75<sup>th</sup> percentile and minimum to maximum distributions of the data. Significant differences from the one-way ANOVA with Duncan's post hoc test are shown for groups: \*\*\*  $p < 0.001$ , \*\*  $p < 0.01$ , \*  $p < 0.05$ , n.s., non significant.



**Figure S13.** Box plots depict (A) carboxylation turnover rate ( $K_{cat}$ :  $s^{-1}$ ), (B) Michaelis-Menten constant for  $CO_2$  at ambient atmospheric  $O_2$  ( $K_C^{21\%O_2}$ :  $\mu M$ ) and (C) specificity of  $CO_2$  to  $O_2$  ( $Sc/o$ :  $mol\ mol^{-1}$ ) predictions made for the wild-type form IB (plants) Rubisco large subunit (RbcL) sequence-space using the fully trained Gaussian process (GP) models with the learned encoding scheme. Predictions are shown for all known land plant Rubiscos. Predictions were grouped by taxonomical type (Angiosperms, Ferns (including Pteridophytes and Lycopodiophytes) Gymnosperms or Bryophytes). Box plot horizontal lines show the median value, and the box and whisker represent the 25<sup>th</sup> and 75<sup>th</sup> percentile and minimum to maximum distributions of the data. Significant differences from the one-way ANOVA with Duncan's post hoc test are shown for groups: \*\*\*  $p < 0.001$ , \*\*  $p < 0.01$ , \*  $p < 0.05$ , n.s., non significant.

**Table S2.** Rubisco experimental kinetics and Rubisco large subunit (RbL) sequences for training Gaussian process models.

Species	Accession No.	Photosynthetic type	Database	Kcat mean	Kcat SD	Sc/o mean	Sc/o SD	Kc <sup>21%O2</sup> mean	Kc <sup>21%O2</sup> SD	Citation
<i>Aegilops biuncialis</i>	LN626633	C <sub>3</sub> plant	EMBL	3.2	0.2	96.3	4.9	16.8	2.9	Prins et al. (2016)
<i>Aegilops comosa</i>	LN626632	C <sub>3</sub> plant	EMBL	2.9	0.2	106.0	4.3	13.5	2.1	Prins et al. (2016)
<i>Aegilops cylindrica</i>	LN626627	C <sub>3</sub> plant	EMBL	3.7	0.2	109.0	3.8	13.7	2.9	Prins et al. (2016)
<i>Aegilops juvenalis</i>	CEH24677.1	C <sub>3</sub> plant	NCBI	3.3	0.2	90.4	6.4	20.6	3.8	Prins et al. (2016)
<i>Aegilops speltoides</i>	LN626629	C <sub>3</sub> plant	EMBL	3.2	0.2	102.0	5.6	16.5	3.2	Prins et al. (2016)
<i>Aegilops tauschii</i>	LN626630	C <sub>3</sub> plant	EMBL	2.9	0.2	107.0	8.3	14.9	1.8	Prins et al. (2016)
<i>Aegilops triuncialis</i>	LN626634	C <sub>3</sub> plant	EMBL	2.6	0.2	103.0	5.9	12.8	2.2	Prins et al. (2016)
<i>Aegilops uniaristata</i>	LN626631	C <sub>3</sub> plant	EMBL	2.7	0.2	102.0	1.1	13.8	2.3	Prins et al. (2016)
<i>Aegilops vavilovii</i>	LN626635	C <sub>3</sub> plant	EMBL	3.3	0.2	109.0	3.0	13.3	1.9	Prins et al. (2016)
<i>Agriophyllum squarrosum</i>	A0A1C3HPL7	C <sub>3</sub> plant	uni	2.8	0.1	93.9	1.1	15.4	1.5	Orr et al. (2016)
<i>Agrostis scabra</i>	A0A1C3HPM5	C <sub>3</sub> plant	uni	3.6	0.1	103.0	1.6	22.0	1.7	Orr et al. (2016)
<i>Agrostis stolonifera</i>	ABK79588.1	C <sub>3</sub> plant	NCBI	5.2	0.1	105.0	1.4	25.3	1.5	Orr et al. (2016)
<i>Amaranthus hybridus</i>	YP_010028517.1	C <sub>4</sub> plant	NCBI	3.8	0.3	82.0	4.0	21.3	1.5	Savir et al. (2010)
<i>Amphicarpaea bracteata</i>	A0A1C3HPM0	C <sub>3</sub> plant	uni	4.0	0.1	97.5	2.2	29.0	2.1	Orr et al. (2016)
<i>Arabidopsis thaliana</i>	QDK58872.1	C <sub>3</sub> plant	NCBI	4.1	0.3	105.0	7.3	16.1	0.7	Galmés et al. (2014b)
<i>Arctagrostis latifolia</i>	A0A1C3HPM4	C <sub>3</sub> plant	uni	5.8	0.1	105.0	1.4	21.0	1.1	Orr et al. (2016)
<i>Artemisia myriantha</i>	A0A1C3HPM2	C <sub>3</sub> plant	uni	3.1	0.1	110.0	1.7	26.4	1.3	Orr et al. (2016)
<i>Artemisia vulgaris</i>	LT576797	C <sub>3</sub> plant	EMBL	3.9	0.2	105.0	2.6	31.9	4.3	Orr et al. (2016)
<i>Avena sativa</i>	AAA84028.1	C <sub>3</sub> plant	NCBI	2.3	0.3	99.9	3.0	18.1	2.0	Hermida-Carrera et al. (2016)
<i>Beta maritima ssp. maritima</i>	AEK33913.1	C <sub>3</sub> plant	NCBI			92.9	3.2			Galmes et al. (2005)
<i>Beta vulgaris</i>	AAY68360.1	C <sub>3</sub> plant	NCBI	2.0	0.3	101.0	2.0	18.6	1.1	Hermida-Carrera et al. (2016)
<i>Brachypodium distachyon</i>	LN626640	C <sub>3</sub> plant	EMBL	2.1	0.1	111.0	4.0	11.9	2.5	Prins et al. (2016)
<i>Brassica oleracea</i>	YP_009564583.1	C <sub>3</sub> plant	NCBI	2.1	0.3	96.2	1.3	19.2	0.3	Hermida-Carrera et al. (2016)
<i>Bromus anomalus</i>	A0A1C3HPM1	C <sub>3</sub> plant	uni	2.9	0.2	101.0	3.2	16.9	3.1	Orr et al. (2016)

<i>Calamagrostis arundinacea</i>	A0A1C3HPL9	C <sub>3</sub> plant	uni	4.1	0.1	104.0	1.6	22.7	1.3	Orr et al. (2016)
<i>Calamagrostis canescens</i>	A0A1C3HPN2	C <sub>3</sub> plant	uni	2.5	0.1	98.5	2.6	15.2	2.2	Orr et al. (2016)
<i>Calamagrostis foliosa</i>	A0A1C3HPN4	C <sub>3</sub> plant	uni	3.5	0.1	105.0	1.4	20.9	1.1	Orr et al. (2016)
<i>Calamagrostis inexpansa</i>	A0A1C3HPM7	C <sub>3</sub> plant	uni	3.3	0.1	111.0	1.3	18.8	1.7	Orr et al. (2016)
<i>Calamagrostis nutkaensis</i>	A0A1C3HPN0	C <sub>3</sub> plant	uni	3.1	0.1	109.0	1.2	20.1	2.8	Orr et al. (2016)
<i>Capsicum annuum</i>	A0A1U8QD66	C <sub>3</sub> plant	uni	1.9	0.1	96.0	4.5	19.8	1.5	Hermida-Carrera et al. (2016)
<i>Cenchrus ciliaris</i>	YP_009573569.1	C <sub>4</sub> plant	NCBI	6.0	0.8	69.9	3.0	29.2	2.3	Sharwood et al. (2016)
<i>Chenopodium album</i>	ATL76552.1	C <sub>3</sub> plant	NCBI	2.9	0.1	78.7	1.0	11.2	2.8	Kubien et al. (2008)
<i>Chenopodium murale</i>	SBO07486.1	C <sub>3</sub> plant	NCBI	4.4	0.2	109.0	0.8	23.8	3.1	Orr et al. (2016)
<i>Chenopodium rubrum</i>	A0A1C3HPN3	C <sub>3</sub> plant	uni	4.1	0.1	97.8	1.0	14.5	1.3	Orr et al. (2016)
<i>Chrysanthellum indicum</i> var. <i>afroamericanum</i>	A0A1C3HPM6	C <sub>4</sub> plant	uni	4.7	0.1	82.4	0.6	28.1	2.3	Orr et al. (2016)
<i>Citrullus ecirrhosus</i>	A0A1C3HPM8	C <sub>3</sub> plant	uni	3.1	0.2	99.9	1.5	18.9	3.0	Orr et al. (2016)
<i>Citrullus lanatus</i> var. <i>lanatus</i>	SBO07491.1	C <sub>3</sub> plant	NCBI	2.5	0.1	107.0	2.4	19.4	1.2	Orr et al. (2016)
<i>Coffea arabica</i>	YP_817490.1	C <sub>3</sub> plant	NCBI	2.1	0.2	98.7	3.8	22.9	2.4	Hermida-Carrera et al. (2016)
<i>Crithmum maritimum</i>	YP_004733335.1	C <sub>3</sub> plant	NCBI	3.4	0.1			18.7	0.2	Galmés et al. (2014b)
<i>Cucurbita maxima</i>	YP_009447458.1	C <sub>3</sub> plant	NCBI	2.2	0.2	98.4	0.4	19.2	1.0	Hermida-Carrera et al. (2016)
<i>Cynodon dactylon</i>	QYC94651.1	C <sub>4</sub> plant	NCBI			89.2	9.0	32.0	2.0	Carmo-Silva et al. (2010)
<i>Dactylis glomerata</i> cv. <i>Porto</i>	AJV89540.1	C <sub>3</sub> plant	NCBI	3.2	0.2			15.7	0.1	Galmés et al. (2014b)
<i>Deschampsia danthonioides</i>	A0A1C3HPP5	C <sub>3</sub> plant	uni	4.5	0.1	108.0	1.6	22.3	1.4	Orr et al. (2016)
<i>Desmodium cinereum</i>	A0A1C3HPP3	C <sub>3</sub> plant	uni	3.0	0.1	97.5	2.0	12.8	1.0	Orr et al. (2016)
<i>Desmodium intortum</i>	A0A1C3HPN7	C <sub>3</sub> plant	uni	3.3	0.1	98.7	1.9	14.2	1.7	Orr et al. (2016)
<i>Desmodium psilocarpum</i>	A0A1C3HPP4	C <sub>3</sub> plant	uni	3.6	0.1	95.9	1.7	15.6	0.9	Orr et al. (2016)
<i>Echinochloa crus-galli</i>	QPB15306.1	C <sub>4</sub> plant	NCBI			83.0	3.0			Jordan and Ogren (1983)

<i>Elymus farctus</i>	A0A1C3HPN9	C <sub>3</sub> plant	uni	3.3	0.1	106.0	1.3	19.5	2.7	Orr et al. (2016)
<i>Eragrostis tef</i>	A0A6C0SV93	C <sub>4</sub> plant	uni	7.1	0.2	89.0	2.1	34.9	2.1	Orr et al. (2016)
<i>Erythrina flabelliformis</i>	A0A1C3HPP1	C <sub>3</sub> plant	uni	3.6	0.1	96.4	1.9	18.4	1.9	Orr et al. (2016)
<i>Espeletia schultzei</i>	AIL52300.1	C <sub>3</sub> plant	NCBI					21.7	2.0	Castrillo (1995)
<i>Euphorbia helioscopia</i>	A0A6M4AEZ2	C <sub>3</sub> plant	uni	1.9	0.1	96.8	1.9	11.5	1.7	Orr et al. (2016)
<i>Euphorbia microsphaera</i>	A0A1C3HPP0	C <sub>3</sub> plant	uni	4.5	0.2	99.7	0.7	25.8	2.7	Orr et al. (2016)
<i>Festuca gigantea</i>	A0A1C3HPN8	C <sub>3</sub> plant	uni	5.1	0.1	108.0	2.4	31.2	1.6	Orr et al. (2016)
<i>Festuca pratensis</i>	K4PCJ4	C <sub>3</sub> plant	uni	5.1	0.2	106.0	1.1	23.1	2.5	Orr et al. (2016)
<i>Flaveria angustifolia</i>	ADW80651.1	C <sub>3</sub> -C <sub>4</sub> plant	NCBI	2.9	0.2	83.2	1.0	13.1	0.1	Kubien et al. (2008)
<i>Flaveria anomala</i>	ADW80650.1	C <sub>3</sub> -C <sub>4</sub> plant	NCBI	3.8	0.2	77.9	0.9	10.7	0.6	Kubien et al. (2008)
<i>Flaveria australasica</i>	P19161.1	C <sub>4</sub> plant	NCBI	3.8	0.0	77.2	0.3	22.0	4.7	Kubien et al. (2008)
<i>Flaveria bidentis</i>	P19161.1	C <sub>4</sub> plant	NCBI	4.2	0.3	75.5	0.8	20.2	1.6	Kubien et al. (2008)
<i>Flaveria chloraefolia</i>	ADW80653.1	C <sub>3</sub> -C <sub>4</sub> plant	NCBI	3.4	0.1	81.6	1.3	12.4	2.3	Kubien et al. (2008)
<i>Flaveria cronquistii</i>	ADW80654.1	C <sub>3</sub> plant	NCBI	3.1	0.3	81.0	2.0	10.8	0.4	Kubien et al. (2008)
<i>Flaveria floridana</i>	ADW80655.1	C <sub>3</sub> -C <sub>4</sub> plant	NCBI	3.2	0.1	84.5	1.1	13.2	0.4	Kubien et al. (2008)
<i>Flaveria kochiana</i>	ADW80656.1	C <sub>4</sub> plant	NCBI	3.7	0.1	77.0	1.2	22.7	2.7	Kubien et al. (2008)
<i>Flaveria linearis</i>	ADW80657.1	C <sub>3</sub> -C <sub>4</sub> plant	NCBI	3.4	0.1	78.1	1.3	12.5	0.9	Kubien et al. (2008)
<i>Flaveria palmeri</i>	ADW80658.1	C <sub>4</sub> plant	NCBI	3.5	0.2	83.8	1.0	13.5	0.9	Kubien et al. (2008)
<i>Flaveria pringlei</i>	ADW80648.1	C <sub>3</sub> plant	NCBI	3.1	0.2	80.8	1.2	12.0	2.1	Kubien et al. (2008)
<i>Flaveria ramosissima</i>	ADW80659.1	C <sub>3</sub> -C <sub>4</sub> plant	NCBI	2.8	0.1	79.8	1.1	12.0	0.2	Kubien et al. (2008)
<i>Flaveria sonorensis</i>	ADW80660.1	C <sub>3</sub> -C <sub>4</sub> plant	NCBI	2.7	0.1	84.3	1.4	10.2	1.4	Kubien et al. (2008)
<i>Flaveria trinervia</i>	ADW80661.1	C <sub>4</sub> plant	NCBI	4.4	0.2	77.0	0.8	17.9	1.8	Kubien et al. (2008)
<i>Flaveria vaginata</i>	ADW80662.1	C <sub>4</sub> plant	NCBI	3.8	0.5	78.7	1.5	21.4	1.9	Kubien et al. (2008)
<i>Flueggea suffruticosa</i>	AFU96118.1	C <sub>3</sub> plant	NCBI	3.4	0.1	101.0	1.5	19.2	1.5	Orr et al. (2016)
<i>Foeniculum vulgare</i>	AMD83923.1	C <sub>3</sub> plant	NCBI	4.4	0.2	94.3	2.7	20.7	2.7	Orr et al. (2016)
<i>Glycine canescens</i>	YP_008145856.1	C <sub>3</sub> plant	NCBI	2.6	0.0	97.1	1.7	17.2	0.9	Orr et al. (2016)
<i>Glycine max</i>	ABC25107.1	C <sub>3</sub> plant	NCBI	1.5	0.1	97.0	1.1	16.2	0.7	Hermida-Carrera et al. (2016)
<i>Helianthus annuus L.</i>	XP_035835866.1	C <sub>3</sub> plant	NCBI			73.6	18.3			Viil et al. (2012)
<i>Hordeum brachyantherum</i>	A0A1C3HPQ0	C <sub>3</sub> plant	uni	2.9	0.1	101.0	2.0	16.2	1.5	Orr et al. (2016)
<i>Hordeum murinum</i>	A0A1C3HPQ4	C <sub>3</sub> plant	uni	4.2	0.1	100.0	1.1	21.5	1.1	Orr et al. (2016)

<i>Hordeum vulgare ssp. vulgare cv. Morex</i>	ASD42799.1	C <sub>3</sub> plant	NCBI					14.9	1.6	Hermida-Carrera et al. (2016)
<i>Ipomoea batatas</i>	AFU50388.1	C <sub>3</sub> plant	NCBI	2.5	0.1	98.5	6.6	21.1	1.0	Hermida-Carrera et al. (2016)
<i>Iris douglasiana</i>	BAO57024.1	C <sub>3</sub> plant	NCBI	3.5	0.2			14.6	0.9	Galmés et al. (2014b)
<i>Lablab purpureus</i>	YP_010046451.1	C <sub>3</sub> plant	NCBI	5.3	0.1	91.1	1.7	21.7	1.6	Orr et al. (2016)
<i>Lactuca sativa</i>	YP_398337.1	C <sub>3</sub> plant	NCBI			94.0	1.9	18.2	1.4	Hermida-Carrera et al. (2016)
<i>Lepidium campestre</i>	A0A1C3HPQ1	C <sub>3</sub> plant	uni	3.4	0.1	92.8	1.2	15.8	1.0	Orr et al. (2016)
<i>Limonium antonii-llorensii</i>	AID50137.1	C <sub>3</sub> plant	NCBI	2.4	0.1	112.0	1.8	13.3	0.5	Galmés et al. (2014a)
<i>Limonium artruchium</i>	AID50126.1	C <sub>3</sub> plant	NCBI	3.0	0.1	113.0	2.0	15.5	0.5	Galmés et al. (2014a)
<i>Limonium balearicum</i>	AID50147.1	C <sub>3</sub> plant	NCBI	3.6	0.1	103.0	1.5	15.6	0.8	Galmés et al. (2014a)
<i>Limonium barceloi</i>	AID50128.1	C <sub>3</sub> plant	NCBI	3.4	0.1	106.0	2.0	14.9	1.0	Galmés et al. (2014a)
<i>Limonium biflorum</i>	AID50121.1	C <sub>3</sub> plant	NCBI	2.4	0.1	112.0	1.8	13.3	0.5	Galmés et al. (2014a)
<i>Limonium companyonis</i>	AID50113.1	C <sub>3</sub> plant	NCBI	3.3	0.2	112.0	2.1	13.3	0.4	Galmés et al. (2014a)
<i>Limonium echioides</i>	AID50149.1	C <sub>3</sub> plant	NCBI	3.9	0.1	106.0	1.8	16.0	0.3	Galmés et al. (2014a)
<i>Limonium ejulabilis</i>	AID50114.1	C <sub>3</sub> plant	NCBI	2.0	0.2	116.0	3.2	11.4	0.3	Galmés et al. (2014a)
<i>Limonium gibertii</i>	AID50136.1	C <sub>3</sub> plant	NCBI	2.5	0.1	112.0	2.5	13.5	1.0	Galmés et al. (2014a)
<i>Limonium grosii</i>	AID50112.1	C <sub>3</sub> plant	NCBI	2.9	0.1	113.0	3.3	13.3	0.7	Galmés et al. (2014a)
<i>Limonium gymnesicum</i>	AID50143.1	C <sub>3</sub> plant	NCBI	2.4	0.1	121.0	2.3	12.6	0.2	Galmés et al. (2014a)
<i>Limonium latebracteatum</i>	BAO57039.1	C <sub>3</sub> plant	NCBI	2.7	0.1			14.2	1.6	Galmés et al. (2014b)
<i>Limonium leonardi-llorensii</i>	AID50109.1	C <sub>3</sub> plant	NCBI	2.8	0.1	110.0	1.1	13.0	0.6	Galmés et al. (2014a)
<i>Limonium magallufianum</i>	AID50120.1	C <sub>3</sub> plant	NCBI	2.6	0.1	111.0	2.9	11.9	0.5	Galmés et al. (2014a)
<i>Limonium retusum</i>	AID50105.1	C <sub>3</sub> plant	NCBI	2.1	0.1	121.0	1.4	10.9	0.6	Galmés et al. (2014a)
<i>Limonium stenophyllum</i>	BAO57038.1	C <sub>3</sub> plant	NCBI	2.6	0.1			12.3	0.3	Galmés et al. (2014b)
<i>Limonium virgatum</i>	AID50151.1	C <sub>3</sub> plant	NCBI	2.4	0.1			13.2	0.3	Galmés et al. (2014b)
<i>Lolium multiflorum</i>	AFV62913.1	C <sub>3</sub> plant	NCBI	4.5	0.1	99.9	1.8	29.1	2.1	Orr et al. (2016)
<i>Lolium perenne</i>	CAO85984.1	C <sub>3</sub> plant	NCBI			80.0	1.0	22.7	1.1	Savir et al. (2010)
<i>Lolium rigidum</i>	A0A1C3HPR5	C <sub>3</sub> plant	uni	4.7	0.1	100.0	1.4	25.0	1.7	Orr et al. (2016)

<i>Macrotyloma uniflorum</i>	ACJ38561.1	C <sub>3</sub> plant	NCBI	4.4	0.2	101.0	2.4	25.2	3.3	Orr et al. (2016)
<i>Manihot esculenta</i>	QFO47553.1	C <sub>3</sub> plant	NCBI	1.4	0.1	101.0	0.9	10.8	0.6	Hermida-Carrera et al. (2016)
<i>Medicago sativa</i>	AOA1B3T043	C <sub>3</sub> plant	uni	1.7	0.1	95.6	2.2	16.4	1.9	Hermida-Carrera et al. (2016)
<i>Megathyrsus maximus</i>	YP_009260243.1	C <sub>4</sub> plant	NCBI	5.3	0.5	80.3	2.8	27.1	1.1	Sharwood et al. (2016)
<i>Mentha aquatica</i>	ADU79049.1	C <sub>3</sub> plant	NCBI			97.2	3.3			Galmes et al. (2005)
<i>Mercurialis annua</i>	AOA1C3HPR2	C <sub>3</sub> plant	uni	3.4	0.2	95.7	2.7	17.0	3.7	Orr et al. (2016)
<i>Musa velutina</i>	UCC34423.1	C <sub>3</sub> plant	NCBI	3.2	0.1	111.0	1.8	19.0	1.9	Orr et al. (2016)
<i>Nicotiana tabacum</i>	1RLC_L	C <sub>3</sub> plant	NCBI	3.1	0.3	80.0	2.6	18.3	0.1	Long et al. (2018)
<i>Oryza barthii/glaberrima</i>	LT576837	C <sub>3</sub> plant	EMBL	2.5	0.0	107.0	1.5	14.0	0.8	Orr et al. (2016)
<i>Oryza eichingeri</i>	LT576838	C <sub>3</sub> plant	EMBL	2.5	0.1	107.0	1.2	14.1	1.5	Orr et al. (2016)
<i>Oryza glaberrima (WAB 1939)</i>	LT576839	C <sub>3</sub> plant	EMBL	2.7	0.0	105.0	1.1	14.9	0.6	Orr et al. (2016)
<i>Oryza glumaepatula</i>	LT576840	C <sub>3</sub> plant	EMBL	2.4	0.1	109.0	0.7	15.2	1.5	Orr et al. (2016)
<i>Oryza longistaminata</i>	LT576841	C <sub>3</sub> plant	EMBL	2.2	0.0	108.0	1.3	15.1	1.1	Orr et al. (2016)
<i>Oryza meridionalis</i>	LT576842	C <sub>3</sub> plant	EMBL	2.6	0.0	107.0	1.3	14.6	0.9	Orr et al. (2016)
<i>Oryza nivara</i>	LT576843	C <sub>3</sub> plant	EMBL	2.7	0.0	107.0	0.6	15.6	1.0	Orr et al. (2016)
<i>Oryza punctata</i>	LT576844	C <sub>3</sub> plant	EMBL	2.7	0.1	109.0	1.2	14.9	2.2	Orr et al. (2016)
<i>Oryza sativa</i>	ANG44656.1	C <sub>3</sub> plant	NCBI	2.1	0.3	93.1	1.2	17.3	2.4	Hermida-Carrera et al. (2016)
<i>Oryza sativa subsp. Indica</i>	LT576845	C <sub>3</sub> plant	EMBL	2.2	0.1	101.0	2.0	7.0	1.6	Orr et al. (2016)
<i>Oryza sativa subsp. Japonica</i>	LT576846	C <sub>3</sub> plant	EMBL	2.8	0.1	106.0	1.4	11.8	1.8	Orr et al. (2016)
<i>Pallenis maritima</i>	BAO57025.1	C <sub>3</sub> plant	NCBI	2.7	0.1			10.6	0.5	Galmés et al. (2014b)
<i>Panicum amarum</i>	LT576849	C <sub>4</sub> plant	EMBL	3.2	0.1	89.5	3.0	33.1	3.9	Orr et al. (2016)
<i>Panicum antidotale</i>	SCM15150.1	C <sub>4</sub> plant	NCBI	3.9	0.2	74.5	0.4			Sharwood et al. (2016)
<i>Panicum bisulcatum</i>	SCM15147.1	C <sub>3</sub> plant	NCBI	2.6	0.4	87.7	1.5	12.6	1.0	Sharwood et al. (2016)
<i>Panicum coloratum</i>	SCM15156.1	C <sub>4</sub> plant	NCBI	3.4	0.6	84.8	2.8	17.3	1.7	Sharwood et al. (2016)
<i>Panicum deustum</i>	SCM15159.1	C <sub>4</sub> plant	NCBI	5.0	0.5	84.8	0.2	28.1	2.0	Sharwood et al. (2016)
<i>Panicum dichotomiflorum</i>	LT576847	C <sub>4</sub> plant	EMBL	3.1	0.1	92.6	1.8	36.3	2.9	Orr et al. (2016)

<i>Panicum milioides</i>	SCM15149.1	C <sub>3</sub> plant	NCBI	2.2	0.3	92.3	1.0	12.1	0.8	Sharwood et al. (2016)
<i>Panicum milliaceum</i>	SCM15155.1	C <sub>4</sub> plant	NCBI			79.9	4.3	13.1	1.1	Sharwood et al. (2016)
<i>Panicum monticola</i>	SCM15151.1	C <sub>4</sub> plant	NCBI	5.3	0.7	79.4	1.7	26.6	4.0	Sharwood et al. (2016)
<i>Panicum phragmitoides</i>	LT576848	C <sub>4</sub> plant	EMBL	2.8	0.3	107.0	1.3	25.1	7.5	Orr et al. (2016)
<i>Panicum virgatum</i>	SCM15154.1	C <sub>4</sub> plant	NCBI	3.3	0.9	82.6	2.8	24.5	2.1	Sharwood et al. (2016)
<i>Paspalum dilatatum</i>	YP_009267237.1	C <sub>4</sub> plant	NCBI			88.0	7.1	30.0	1.2	Carmo-Silva et al. (2010)
<i>Petroselinum crispum</i>	YP_004733851.1	C <sub>3</sub> plant	NCBI			77.0	2.0			Jordan and Ogren (1983)
<i>Phaseolus carteri</i>	LT576850	C <sub>3</sub> plant	EMBL	3.2	0.1	84.5	2.1	14.2	1.9	Orr et al. (2016)
<i>Phaseolus coccineus</i>	LT576851	C <sub>3</sub> plant	EMBL	3.9	0.2	104.0	2.9	15.6	2.6	Orr et al. (2016)
<i>Phaseolus lunatus</i>	LT576852	C <sub>3</sub> plant	EMBL	3.2	0.1	101.0	0.7	17.3	1.1	Orr et al. (2016)
<i>Phaseolus vulgaris</i>	QCQ20610.1	C <sub>3</sub> plant	NCBI	1.7	0.2	99.7	2.7	14.0	1.0	Hermida-Carrera et al. (2016)
<i>Pisum sativum</i>	AIK21409.1	C <sub>3</sub> plant	NCBI			90.2	1.7			Uemura et al. (1997)
<i>Plantago lanceolata L.</i>	AAA84554.1	C <sub>3</sub> plant	NCBI			77.3	12.9			Viil et al. (2012)
<i>Poa palustris</i>	LT576854	C <sub>3</sub> plant	EMBL	4.2	0.1	112.0	2.1	19.2	1.3	Orr et al. (2016)
<i>Portulaca oleracea</i>	AFB70630.1	C <sub>4</sub> plant	NCBI	5.9	0.4	78.0	4.0			Savir et al. (2010)
<i>Puccinellia distans</i>	LT576855	C <sub>3</sub> plant	EMBL	5.4	0.2	104.0	1.9	22.2	2.4	Orr et al. (2016)
<i>Puccinellia lemmonii</i>	LT576856	C <sub>3</sub> plant	EMBL	5.2	0.4	102.0	1.0	28.1	5.9	Orr et al. (2016)
<i>Puccinellia maritima</i>	LT576857	C <sub>3</sub> plant	EMBL	5.4	0.2	106.0	1.9	20.8	2.4	Orr et al. (2016)
<i>Puccinellia nuttalliana</i>	LT576858	C <sub>3</sub> plant	EMBL	4.0	0.1	105.0	1.1	25.2	1.8	Orr et al. (2016)
<i>Pueraria montana</i>	QRM91166.1	C <sub>3</sub> plant	NCBI	2.7	0.1	101.0	1.0	20.9	1.6	Orr et al. (2016)
<i>Saccharum officinarum</i>	ASV52260.1	C <sub>4</sub> plant	NCBI	3.9	0.3	82.2	1.8	31.7	2.1	Hermida-Carrera et al. (2016)
<i>Secale cereale</i>	LN626639	C <sub>3</sub> plant	EMBL	3.2	0.2	91.5	4.1	20.2	2.1	Prins et al. (2016)
<i>Setaria italica</i>	AGZ13153.1	C <sub>4</sub> plant	NCBI			58.0	2.0			Jordan and Ogren (1983)
<i>Setaria viridis</i>	AGT56139.1	C <sub>4</sub> plant	NCBI	5.9	0.5	72.7	0.2	25.5	2.0	Sharwood et al. (2016)
<i>Sideritis cretica subsp. spicata</i>	BAO57028.1	C <sub>3</sub> plant	NCBI	2.0	0.3			12.8	0.2	Galmés et al. (2014b)
<i>Solanum lycopersicum</i>	ANW81480.1	C <sub>3</sub> plant	NCBI	2.3	0.2	92.4	2.3	16.6	1.4	Hermida-Carrera et al. (2016)
<i>Solanum tuberosum</i>	ABD47065.1	C <sub>3</sub> plant	NCBI	2.0	0.3	95.4	2.3	18.0	0.8	Hermida-Carrera et al. (2016)
<i>Sorghum bicolor</i>	YP_899415.1	C <sub>4</sub> plant	NCBI	5.4	0.1	70.0	1.0			Savir et al. (2010)



<i>Sphenostylis stenocarpa</i>	LT576860	C <sub>3</sub> plant	EMBL	2.8	0.1	93.6	3.4	17.4	1.5	Orr et al. (2016)
<i>Spinacia oleracea</i>	CAA23473.1	C <sub>3</sub> plant	NCBI	2.4	0.1	97.0	1.2	26.9	0.8	Hermida-Carrera et al. (2016)
<i>Steinchisma laxa</i>	YP_009261073.1	C <sub>3</sub> plant	NCBI	2.3	0.3	91.4	4.8			Sharwood et al. (2016)
<i>Tephrosia candida</i>	LT576861	C <sub>3</sub> plant	EMBL	2.2	0.2	97.8	2.5	15.9	3.8	Orr et al. (2016)
<i>Tephrosia purpurea</i>	LT576862	C <sub>3</sub> plant	EMBL	2.2	0.1	103.0	1.1	12.4	1.4	Orr et al. (2016)
<i>Tephrosia rhodesica</i>	LT576863	C <sub>3</sub> plant	EMBL	2.2	0.1	91.6	3.3	13.7	1.2	Orr et al. (2016)
<i>Tetragonia expansa</i>	YP_009462736.1	C <sub>3</sub> plant	NCBI			81.0	1.0	17.6	1.2	Savir et al. (2010)
<i>Teucrium heterophyllum</i>	BAO57027.1	C <sub>3</sub> plant	NCBI	2.7	0.1			10.6	0.2	Galmés et al. (2014b)
<i>Trachycarpus fortunei</i>	YP_010010626.1	C <sub>3</sub> plant	NCBI	2.8	0.1			14.2	1.3	Galmés et al. (2014b)
<i>Trifolium repens</i>	QGV12450.1	C <sub>3</sub> plant	NCBI					20.8	1.2	Lehnher et al. (1985)
<i>Triticale</i>	LN626638	C <sub>3</sub> plant	EMBL	3.6	0.2	99.7	5.1	16.1	2.4	Prins et al. (2016)
<i>Triticum aestivum</i>	BCU02136.1	C <sub>3</sub> plant	NCBI	2.2	0.2	100.0	1.8	16.0	0.6	Hermida-Carrera et al. (2016)
<i>Triticum aestivum SATYN1</i>	LN626617	C <sub>3</sub> plant	EMBL	3.3	0.2	93.8	3.2	17.6	2.5	Prins et al. (2016)
<i>Triticum aestivum SATYN2</i>	LN626618	C <sub>3</sub> plant	EMBL	3.3	0.2	96.2	3.8	17.9	2.8	Prins et al. (2016)
<i>Triticum aestivum SATYN3</i>	LN626619	C <sub>3</sub> plant	EMBL	3.6	0.2	92.9	5.9	15.7	2.2	Prins et al. (2016)
<i>Triticum dicoccon1</i>	LN626623	C <sub>3</sub> plant	EMBL	3.6	0.2	94.1	3.8	15.4	2.1	Prins et al. (2016)
<i>Triticum dicoccon2</i>	LN626624	C <sub>3</sub> plant	EMBL	3.6	0.2	90.9	1.7	13.5	2.3	Prins et al. (2016)
<i>Triticum dicoccon3</i>	LN626625	C <sub>3</sub> plant	EMBL	3.4	0.2	95.2	4.2	16.8	2.5	Prins et al. (2016)
<i>Triticum dicoccon4</i>	LN626626	C <sub>3</sub> plant	EMBL	3.6	0.2	93.8	3.2	18.6	2.7	Prins et al. (2016)
<i>Triticum monococcum</i>	LN626620	C <sub>3</sub> plant	EMBL	3.2	0.2	104.0	4.1	14.0	2.1	Prins et al. (2016)
<i>Triticum timonovum</i>	LN626622	C <sub>3</sub> plant	EMBL	3.5	0.2	101.0	3.2	16.8	3.3	Prins et al. (2016)
<i>Triticum timopheevii</i>	LN626621	C <sub>3</sub> plant	EMBL	3.4	0.2	102.0	2.8	16.2	2.3	Prins et al. (2016)
<i>Urochloa mosambicensis</i>	SCM15160.1	C <sub>4</sub> plant	NCBI	5.7	0.7	82.5	1.3			Sharwood et al. (2016)
<i>Urochloa panicoides</i>	SCM15158.1	C <sub>4</sub> plant	NCBI	5.6	0.6	78.3	0.3	24.1	1.0	Sharwood et al. (2016)
<i>Zea mays cv. Carella</i>	CAA78027.1	C <sub>4</sub> plant	NCBI	4.1	0.6	87.3	1.4	42.0	2.8	Hermida-Carrera et al. (2016)
<i>Zostera japonica</i>	BAD05105.1	C <sub>4</sub> plant	NCBI			84.1	7.7	28.1	1.8	Carmo-Silva et al. (2010)

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